SEQUENCE LISTING

<110> HANAGATA, HIROSHI NISHIJYO, TAKAYUKI
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<140> 10/578,781 <141> 2006-05-09
<150> PCT/JP04/16912 <151> 2004-11-08
<150> JP 2003-381606 <151> 2003-11-11
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gat gtg aga ttt tcg acc tat gcg gtg cca atg atc atc gga gaa att 288 Asp Val Arg Phe Ser Thr Tyr Ala Val Pro Met Ile Ile Gly Glu Ile 85 90 95

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	gaa Glu															384
	ttc Phe 130			-				-					_			432
	acg Thr		-			_					-	_				480
	tcc Ser					_		_		_		_				528
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	gcc Ala	_		_	_				_	_		_		_		624
	gtc Val 210			-			_		-		-			-	_	672
-	cgt Arg				_		_		_		-		_	_	_	720
	ctg Leu		_		_				_		tag					756
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Gln	Val	Lys	Asp	Leu	Ile	Ala	Lys	Ser	Gln	Ala	Gly	Asp	Thr	Asp	Ala	

Gln Val Lys Asp Leu Ile Ala Lys Ser Gln Ala Gly Asp Thr Asp Ala 25 Gln Gln Ala Gly Asp Thr Asp Ala 30 Arg Glu Leu Leu Val Asn Ser Asn Ale 40 Leu Val Trp Ser Val Val Gln Arg Phe Ile Asn Arg Gly Tyr Glu Ala Asp Asp Leu Phe Gln Ile 50

Gly 65	Cys	Ile	Gly	Leu	Leu 70	Lys	Ala	Val	Asp	Lys 75	Phe	Asp	Leu	Ser	Tyr 80	
Asp	Val	Arg	Phe	Ser 85	Thr	Tyr	Ala	Val	Pro 90	Met	Ile	Ile	Gly	Glu 95	Ile	
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Gln	Phe 130	Gly	Arg	Ala	Pro	Thr 135	Ile	Ala	Glu	Val	Ala 140	Glu	Ala	Val	Gly	
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Ile	Ala	Leu 195	Lys	Asp	Ala	Ile	Ser 200	Arg	Leu	Ser	Glu	Arg 205	Glu	Gln	Leu	
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	att Ile 530															1632
	ggc Gly															1680
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	cgc Arg 610															1872
	tac Tyr															1920
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cgt Arg	tac Tyr	tgg Trp	aat Asn 660	aac Asn	gat Asp	gag Glu	ggc Gly	aac Asn 665	gag Glu	gag Glu	gcc Ala	att Ile	gcc Ala 670	gac Asp	tcc Ser	2016
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	gat Asp 690															2112

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Lys Lys Ala Ser Lys		Ser Ala Val Ser I	Lys Ala Ala Ser
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Thr Thr Pro Glu Gly		Thr Thr Met Ala I	Lys Tyr Tyr Tyr
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- Gly Asn Asp Ala Asn Pro Arg Asp Leu Val Val Glu Thr Leu Glu Ser 225 230 235 240
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- Leu Asp Asn Leu Met Leu Val His Ser Gly Ile Gly Glu Glu Thr Gly 275 280 285
- Glu Asp Ala Asp Ala Ile Trp Ser His Arg Trp Thr Leu Lys Lys Pro 290 295 300
- Thr Glu Ile Pro Gly Thr Ser Leu Lys Ala Tyr Asp Tyr Met Ile Gln 305 310 315 320
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- Tyr Glu Asp Leu Lys Lys Arg Lys Lys Gln Ala Ser Leu Tyr Asp Gly 405 410 415
- Ser Ser Leu Asp Glu Asp Gly Lys Val Ile Lys Leu Asn Met Pro Gln 420 425 430
- Val Glu Lys Thr Pro Pro Val Gln Pro Lys Asp Gly Asp Tyr Ser Tyr 435 440 445
- Phe Ser Asp Glu Gly Asp Asn Leu Asn Thr Lys Met Thr Ser Glu Val 450 460
- Ile Asp Leu Thr Gly Ala Ser Ser Ala Ser Met Ser Phe Asp Ser Trp 465 470 475 480
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Val Asp Ser Gly Glu Ser Thr Thr Val Lys Glu Tyr Asp Asp Glu Thr 500 505 510

Lys Gly Trp Asp Lys Glu Glu Ile Ser Leu Asn Asp Phe Ala Gly Lys 515 520 525

Lys Ile Gln Val Glu Phe Asn Tyr Val Thr Asp Gly Gly Leu Ala Met 530 540

Ser Gly Phe Tyr Leu Asp Asn Phe Ala Val Thr Ala Asp Gly Glu Val 545 550 560

Val Phe Ser Asp Asp Ala Glu Gly Asp Gln Lys Phe Asp Leu Asp Gly 565 570 575

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Val Glu Leu Arg Ser His Glu Gly Val Asp Glu Gly Leu Lys Tyr Phe 595 600 605

Arg Arg Asn Asp Thr Phe Phe Thr Tyr Asp Pro Gly Leu Val Ile Trp 610 615 620

Tyr Tyr Asp Gly Arg Phe Gly Lys Thr Gln Asp Asn Asn Thr Ser Asn 625 630 635 640

His Pro Gly Tyr Gly Met Leu Gly Val Val Asp Ala His Gl
n Glu Val 645 650 655

Arg Tyr Trp Asn Asp Glu Gly Asn Glu Glu Ala Ile Ala Asp Ser 660 665 670

Arg Tyr Gln Val Asn Asp Ala Ala Phe Ser Pro Asn Lys Thr Ser Gly 675 680 685

Met Asp Leu Asp Tyr Ile Leu Gly Thr Met Asp Tyr Glu Pro Leu Lys 690 695 700

Gly Ile Thr Val Phe Lys Asp Ser Asp Asp Tyr Thr Met Pro Glu Val 705 710 715 720

Pro Glu Ile Gly Lys Ile Leu Pro Lys Ile Gly Leu Gln Ile Lys Leu
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	ctg Leu															768
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Tyr Gln Ser Leu Tyr Gly Arg Leu Gln Glu Val Ala Glu Thr Leu Pro \$35\$ \$40\$ \$45\$

Pro Leu Tyr Arg Glu His Val Tyr His Pro Phe Leu Gln Ala Met Asp 50 55 60

Lys Leu Ser Glu Ser Gly Phe Ala Gln Met Leu Arg Arg Asp Pro Gln 65 70 75 80

Lys Glu Arg Glu Ala Gly Leu Phe Cys Asp Ile Ala Gln Ala Ile Leu 85 90 95

Gln Asn Gly Glu Ala Tyr Glu Arg Asp Ala Thr Asp Ala Phe Gln Glu 100 105 110

Val Val Ser Asp Leu Tyr Asp Gly Phe Leu Ser Glu Glu Asp Arg Ser 115 120 125

Gly Ile Lys Pro Pro Asp Glu Ser Leu Ile Ala Pro Leu Val Lys Trp 130 135 140

Gly Arg Pro Gln Phe Gly Pro Tyr Thr Trp Thr Ala Glu Ala Ala Ala 145 150 155

His Phe Gly Ile Lys Thr Gly Ile Val Asn Leu Pro Pro Ala Asn Ala 165 170 175

Arg Leu Gly Leu Leu Ala Trp Ser Ala Leu Gly His Glu Thr Ala Gly 180 185

His Asp Ile Leu His Ala Asp Thr Gly Leu Leu Gly Glu Leu Gln Gln
195 200 205

Thr Val Tyr Asp Ala Leu Phe Asp Glu Leu His Asn Arg Thr Leu Ala 210 $\,$ 215 $\,$ 220 $\,$

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Gly Pro Gln Asn Asp Pro His Pro Ala Asp Ile Leu Arg Gly Tyr Leu 275 280 285

Ala Ala Glu Thr Ala Arg Leu Leu His Phe Asp Asn Ala Ser Asp Trp 290 295 300

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Ile Gly Arg Ala Ser Leu Asp Ala Glu Thr Ala Gln Lys Ser Ala Ala 325 330 335

Ile Val Ala Arg Thr Ile Met Glu Ala Arg Leu Leu Ser Leu Glu Gly 340 345 350

His Ala Leu Gly Gln Ile Gln Asn Trp His Asn Glu Asp Glu Arg Ile 355 360 365

Val Gln Glu Ile Arg Ser His Phe Thr Gly Ser Leu Thr Val Gln Asp 370 375 380

Gly Ile Val Ser Gly Met Tyr Ala Ala His Val Val Ala Ala Val 385 390 395 400

Gln Ala Ala Val Ser Gly Glu Met Asp Thr Ser Ala Ala Phe Thr Gly 405 410 415

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